SEQUENCE LISTING

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<110> PAIGE, Lisa A.
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      BUEHRER, Benjamin
<120> METHOD OF PREDICTING THE ABILITY OF COMPOUNDS TO
      MODULATE THE BIOLOGICAL ACTIVITY OF RECEPTORS
<130> PAIGE1D
<140> 09/429,331
<141> 1999-10-28
<150> PCT/US99/06664
<151> 1999-03-26
<150> 60/082,756
<151> 1998-04-23
<150> 60/099,656
<151> 1998-09-09
<150> 60/115,345
<151> 1999-01-08
<160> 360
<170> PatentIn Ver. 2.0
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Gly Ser Gly Lys
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<213> Artificial Sequence

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Ser Ser Trp Asp Met His Gln Phe Phe Trp Glu Gly Val Ser Arg
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Ser Ser Pro Gly Ser Arg Glu Trp Phe Lys Asp Met Leu Ser Arg
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<223> N at each occurrence is A, C, G or T; K at each
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agtgtgtgcc tcgagannkn nknnknnknn knnknnkctg nnknnkctgc tgnnknnknn 60
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Ser Ser His Trp Ser Ser Asp Ser Ile Phe Pro Gly Phe Trp Tyr Ser
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Gly
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Ser Arg Gly Gly Val Asp Leu Asp Ile Gly Asn Ser Ala
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Glu Gly Glu Asp Val Arg Thr Arg Ile Ala Asn
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Ser Ser Trp Val Arg Leu Ser Asp Phe Pro Trp Gly Val Ser Arg
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Ser Ser Trp Asp Arg Leu Ser Asp Phe Pro Trp Gly Val Ser Arg
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Ser Ser Trp Ile Arg Leu Arg Asp Leu Pro Trp Gly Glu Ser Arg
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Ser Ser Gly Ile Cys Phe Phe Trp Asp Gly Cys Phe Glu Ser Arg
<210> 325
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      peptide
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His His His Arg His Pro Ala His Pro His Thr Tyr Gly Gly
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Ser Arg Ala Gly Leu Leu Ser Asp Leu Leu Glu Gly Lys Ser Arg
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Ser Ser Arg Ser Leu Leu Arg Asp Leu Leu Met Val Asp Ser Arg
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<210> 329
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peptide

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<400> 329
Ser Ser Asn Lys Leu Leu Tyr Asn Leu Leu Lys Met Glu Ser Arg
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<212> PRT
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Ser Ser Lys Ser Leu Leu Leu Asn Leu Leu Ser Thr Pro Ser Arg
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His Ser Phe Pro Arg Glu Ser Leu Leu Val Arg Leu Leu Gln Gly Gly
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                                     10
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Ser Arg Leu Glu Glu Leu Leu Lys Trp Gly Ser Val Thr Ser Arg
                                     10
<210> 334
<211> 15
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<213> Artificial Sequence
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     peptide
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Ser Arg Leu Glu Gln Leu Leu Lys Glu Glu Phe Ser Tyr Ser Arg
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<212> PRT
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Ser Arg Leu Glu Gln Leu Leu Arg Ser Glu Pro Asp Phe Ser Arg
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Ser Arg Leu Glu Asp Leu Leu Arg Ala Pro Phe Thr Thr Ser Arg
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Ser Arg Leu Glu Ser Leu Leu Arg Phe Gly Gln Leu Asp Ser Arg
                  5
                                     10
 1
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Ser Arg Leu Glu Glu Leu Leu Gly Thr Asn Arg Asp Ser Arg
                                     10
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                  5
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      peptide
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<400> 343
Ser Arg Leu Ser Cys Leu Leu Met Gly Phe Glu Asp Cys Ser Arg
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Ser Ser Arg Leu Trp Gln Leu Leu Ala Ser Thr Asp Thr Ser Arg
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Ser Arg Ala Gly Pro Val Leu Trp Gly Leu Leu Ser Glu Ser Arg
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<210> 353
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<400> 353
Ser Ser Glu Tyr Cys Phe Tyr Asp Ser Ala His Cys Ser Arg
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Ser Arg Ser Leu Leu Glu Cys His Leu Met Gly Asn Cys Ser Arg
                                      10
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Ser Ser Glu Leu Leu Arg Trp His Leu Thr Arg Asp Thr Ser Arg
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<211> 15
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Ser Arg Leu Glu Tyr Trp Leu Lys Trp Glu Pro Gly Pro Ser Arg
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Ser Arg Ser Asp Ser Ile Leu Trp Arg Met Leu Ser Glu Ser Arg
                  5
<210> 358
<211> 16
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peptide

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Ser Ser Lys Gly Val Leu Trp Arg Met Leu Ala Glu Pro Val Ser Arg
 1 5
                            10
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<223> Description of Artificial Sequence: Arbitrary
    peptide
<400> 359
His Ser His Gly Pro Leu Thr Leu Asn Leu Leu Arg Ser Ser Gly Gly
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Ser Ser Ala Gly Gly Gly Ala Pro Ala Gly Ser Thr Pro Ser Arg
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